NCBI Entrez BLAST 2 sequences BLAST Example Help

# **BLAST 2 SEQUENCES**

This tool produces the alignment of two given sequences using <u>BLAST</u> engine for local alignment. The stand-alone executable for blasting two sequences (bl2seq) can be retrieved from <u>NCBI ftp site</u> <u>Reference:</u> Tatiana A. Tatusova, Thomas L. Madden (1999), "Blast 2 sequences - a new tool for comparing protein and nucleotide sequences", FEMS Microbiol Lett. 174:247-250

Program blastp ▼ Matrix BLOSUM62 ▼			
Parameters used in <u>BLASTN</u> program only:  Reward for a match: Penalty for a mismatch:			
☐ Use Mega BLAST Strand option Not Applicable ▼		, =	
Open gap 11 and extension gap 1 penalties gap x_dropoff 50 expect 10.0 word size 3 Filter Align			
Sequence 1 Enter accession or GI or download from file or sequence in FASTA format from: 0 to: 0			
nqs snf gpm kgg nfg grs sgp ygg ggq yfa kpr nqg gy		,	
Sequence 2 Enter accession or GI or download from file			
or sequence in FASTA format from: 0 to: 0		,	•
YDR RGR PGD GMV GFS ADE TWD SAI DTW SPS EWQ MAY			·
Align Clear Input	<b></b>		· .

Comments and suggestions to <u>blast-help@ncbi.nlm.nih.gov</u>



# **Blast 2 Sequences results**

PubMed Entrez BLAST OMIM Taxonomy Structure

**BLAST 2 SEQUENCES RESULTS VERSION BLASTP 2.2.5 [Nov-16-2002]** 

Matrix BLOSUM62 ▼ gap open: 11 gap extension: 1

x\_dropoff: 50 expect: 10.0 wordsize: 3 Filter ✓ Align

Sequence 1 lcl|seq\_1 Length 38

Sequence 2 |cl|seq\_2 | Length 36

No significant similarity was found

# (FILE 'HOME' ENTERED AT 13:41:15 ON 12 JUN 2003)

#### FILE 'MEDLINE' ENTERED AT 13:45:48 ON 12 JUN 2003

- L1 0 S NON-CLASSICAL NUCLEOAR LOCALIZATION SIGNAL
- L2 1 S NON-CLASSICAL NUCLEAR LOCALIZATION SIGNAL

FILE 'CAPLUS, EMBASE, BIOSIS, BIOTECHDS, SCISEARCH' ENTERED AT 13:47:32

ON 12 JUN 2003

- L3 6 S L2
- L4 2 DUP REM L3 (4 DUPLICATES REMOVED)

## (FILE 'HOME' ENTERED AT 14:31:19 ON 12 JUN 2003)

### FILE 'MEDLINE' ENTERED AT 14:31:43 ON 12 JUN 2003

- L1 0 S STABLY TRANSFECTED AND SCAFFOLD AND NUCLEAR LOCALIZATION
- L2 10 S STABLY TRANSFECTED AND SCAFFOLD
- L3 17 S PLASMID AND EBNA-1 AND EPISOM? AND (STABLE OR STABLY)
- L4 8 S PLASMID AND LARGE T AND EPISOM? AND (STABLE OR STABLY)
- L5 326 S (STABLY OR STABLE) AND EPISOM?
- L6 37 S L5 AND (EBNA OR LARGE T)

=> file stnguide